Att'y Dkt. No. US-111

U.S. App. No: 10/673,860

## IN THE CLAIMS:

Kindly rewrite Claims 1-11 as follows, in accordance with 37 C.F.R. § 1.121:

- 1. [currently amended] A method for producing a heterologous protein which comprises comprising
- <u>bacterium or mutant thereof mutant coryneform bacterium having a genetic expression construct wherein comprising a nucleic acid sequence encoding a signal peptide region derived from a coryneform bacterium which is downstream of a promoter sequence which functions in a coryneform bacterium, and a nucleic acid sequence encoding a heterologous protein which is downstream of said nucleic acid sequence encoding said signal peptide region, said mutant coryneform bacterium having a capacity of secreting the heterologous protein at least 2 fold higher than the wild type Corynebacterium glutamicum ATCC13869, and</u>
- allowing said mutant coryneform bacterium to produce said heterologous
  protein, and
- <u>B3</u>) recovering said heterologous protein, wherein said bacterium or mutant thereof is able to secrete the heterologous protein at least 2-fold higher than *Corynebacterium glutamicum* ATCC13869 having said genetic expression construct.

## 2. [cancelled]

- 3. [currently amended] The method of claim 1, wherein said mutant coryneform bacterium does not produce a cell surface protein and which is derived from Corynebacterium glutamicum AJ12036 (FERM BP-734).
- 4. [previously presented] The method of claim 1, wherein said signal peptide region comprises a signal peptide of a cell surface protein from a coryneform bacterium.
- 5. [withdrawn] The method of claim 1, wherein said signal peptide region comprises a

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signal peptide of a cell surface protein from Corynebacterium glutamicum.

- 6. [withdrawn] The method of claim 1, wherein said signal peptide region comprises the amino acid sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2.
- 7. [currently amended] The method of claim 1, wherein said signal peptide region comprises a signal peptide of a cell surface protein derived-from Corynebacterium ammoniagenes.
- 8. [previously presented] The method of claim 7, wherein said signal peptide comprises the amino acid sequence of SEQ ID NO: 3.
- 9. [withdrawn] The method of claim 5, wherein said signal peptide comprises a sequence having at least one replacement, deletion, addition, or insertion of an amino acid, or a combination thereof in the amino acid sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2.
- 10. [currently amended] The method of claim 1, wherein said culturing of said mutant eoryneform bacterium or said mutant thereof is conducted in a medium containing at least 0.25 g/l (2.25mM) of calcium ion.
- 11. [currently amended] The method of claim 1, wherein said culturing of the said mutant eoryneform—bacterium or said mutant thereof is conducted by controlling the dissolved oxygen concentration at 3% or less.
- 12. [withdrawn] The method of claim 7, wherein said signal peptide comprises a sequence having at least one replacement, deletion, addition, or insertion of an amino acid, or a combination thereof in the amino acid sequence of SEQ ID NO:3.